

GenCore version 5.1.7
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OM nucleic - nucleic search, using sw model

Run on: April 15, 2006, 05:59:36 ; Search time 8424 Seconds
(without alignments)
10911.190 Million cell updates/sec

Title: US-10-796-669A-2

Perfect score: 1617

Sequence: 1 atgactaaaaaccggcacg.....atgaccatttcaaagcttaa 1617

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 5883141 seqs, 28421725653 residues

Total number of hits satisfying chosen parameters: 11766282

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : GenEmbl:
1: gb_ba:
2: gb_in:
3: gb_env:
4: gb_om:
5: gb_ov:
6: gb_pat:
7: gb_ph:
8: gb_pr:
9: gb_ro:
10: gb_sts:
11: gb_sy:
12: gb_un:
13: gb_vi:
14: gb_htg:
15: gb_pl:

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OM nucleic - nucleic search, using sw model

Run on: April 15, 2006, 05:56:21 ; Search time 1038 Seconds
(without alignments)
10382.281 Million cell updates/sec

Title: US-10-796-669A-2

Perfect score: 1617

Sequence: 1 atgactaaaaaccggcacg.....atgaccatttcaaagcttaa 1617

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 4996997 seqs, 3332346308 residues

Total number of hits satisfying chosen parameters: 9993994

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : N_Geneseq_21:
1: geneseqn1980s:
2: geneseqn1990s:
3: geneseqn2000s:
4: geneseqn2001as:
5: geneseqn2001bs:
6: geneseqn2002as:
7: geneseqn2002bs:
8: geneseqn2003as:
9: geneseqn2003bs:
10: geneseqn2003cs:
11: geneseqn2003ds:
12: geneseqn2004as:
13: geneseqn2004bs:
14: geneseqn2005s:

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OM nucleic - nucleic search, using sw model

Run on: April 15, 2006, 06:07:01 ; Search time 6890 Seconds
(without alignments)
10980.365 Million cell updates/sec

Title: US-10-796-669A-2
Perfect score: 1617
Sequence: 1 atgactaaaaaccggcacg.....atgaccatttcaaagcttaa 1617

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 41078325 seqs, 23393541228 residues

Total number of hits satisfying chosen parameters: 82156650

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : EST:
1: gb_est1:
2: gb_est2:
3: gb_est3:
4: gb_htc:
5: gb_est4:
6: gb_est5:
7: gb_est6:
8: gb_est7:
9: gb_gss1:
10: gb_gss2:
11: gb_gss3:

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